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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: Wed Aug 08 11:54:07 EDT 2007

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Application No: 10780638 Version No: 2.0

Input Set:

Output Set:

Started: 2007-08-07 17:13:50.435
Finished: 2007-08-07 17:13:52.249
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 814 ms
Total Warnings: 33
Total Errors: 0
No. of SeqIDs Defined: 33
Actual SeqID Count: 33

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W 213	Artificial or Unknown found in <213> in SEQ ID (19)
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Input Set:

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Error code	Error Description
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SEQUENCE LISTING

<110> Commonwealth Scientific and Industrial Research Organization

<120> Efficient gene silencing in plants using short dsRNA sequences

<130> BCS-03-2001

<140> 10780638

<141> 2004-02-19

<150> US 60/447,711

<151> 2003-02-19

<160> 33

<170> PatentIn version 3.3

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<211> 341

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thaliana var. Landsberg erecta

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<223> SalI restriction site

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<223> poly T nucleotide stretch

<220>

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tctcttcttc tatcttaat ctgtgtatac tatttcacta ttgggcttt tattagtcta 180
taatggact caaaaataagg ctttgccca catcaaaaag ataagtaca aatcaaaaact 240
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<223> poly T stretch

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ctctcttctt ctatctaaa tctgtgtata ctatttcact attgggcttt ttattagtct 180
ataatggac taaaaataag gctttggccc acataaaaaa gataagtac aaatcaaaac 240
taaattcaga gtctttctc ccacatcggt cactgtactc ttttgtgttt gtttatatat 300

tacacgaacc gatcttggt acgtcgagct aagtaacatg agcttgcAAC ccatgtgggg 360
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tttctcgag 429

<210> 3
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<223> PvuI restriction site

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<223> poly T stretch

<220>
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<222> (329)..(334)
<223> EcoRI restriction site

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agatctgttt tggctatgtt ggacgaaaca agtgaacttt taggatcaac tttagtttat 120
atatggagct tatatcgagc aataagataa gtggggcttt tatgtatTTT aatggcgat 180
cgtccataga ttcaactaata cccatgccca gtacccatgt atgcgtttca tataagctcc 240
taatttctcc cacatcgctc aaatctaaac aaatcttggtt gtatataaa cactgaggga 300
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 <223> XhoI restriction site

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 <222> (453)..(461)
 <223> poly T stretch

<220>
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 <222> (462)..(467)
 <223> EcoRI restriction site

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 agatctgttt tggctatgtt ggacgaaaca agtgaacttt taggatcaac tttagtttat 120
 atatggagct tatatcgagc aataagataa gtgggccttt tatgttaattt aatgggctat 180
 cgtccataga ttcactaata cccatgccca gtaccatgt atgcgtttca tataagctcc 240
 taatttctcc cacatcgctc aaatctaaac aaatcttggtt gtatatataa cactgaggga 300
 gcaacattgg tcacgacctt acttgaacag gatctgtct ataggctcgt acctctgttt 360
 ctttgatttc tcaagagaca ggcccttaac cctgggtgat gaaccatgac cgtgcggcta 420
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<220>
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<220>
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 ttagcttttt ttcttcttct tcgttcatac agttttttt tgtttatcg cttacattt 120
 cttgaaccgt agcttcgtt ttcttcttt taacttcca ttggagttt ttgtatctg 180
 ttcatagtt tgtccagga ttagaatgt taggcatacg acctcaaga atttgatgt 240
 ataaaaacatc ttcatctta agatatgaag ataatctca aaaggcccct gggaatctga 300
 aagaagagaa gcaggccat ttatatggaa aagaacaata gtattctta tataggccca 360
 tttaagttga aaacaatctt caaaaagtccc acatcgctta gataagaaaa cgaagctgag 420
 ttatataca gctagagtcg acttttttt gagctc 456

<210> 6
 <211> 488
 <212> DNA
 <213> Artificial Sequence

<220>

<223> sequence of the promoter of the U6-26 snRNA gene of Arabidopsis thaliana var. Landsberg erecta including 20 bases downstream of the transcription initiation site, followed by a unique restriction site in front of an oligo dT stretch

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cttgaaccgt agcttcgtt ttcttctttt taactttcca ttccggagttt ttgtatcttg 180

tttcatagtt tgtcccgaga tttagaatgat taggcatacga accttcaaga atttgattga 240

ataaaaacatc ttcatctta agatatgaag ataatttca aaaggccccct ggaaatctga 300

aagaagagaa gcagggccat ttatatggga aagaacaata gtatttctta tataggcccc 360

tttaagttga aaacaatctt caaaagtccc acatcgctta gataagaaaa cgaagctgag 420

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<220>

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<223> Pol III promoter region

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<221> misc_feature

<222> (385)..(390)

<223> PvuI restriction site

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<222> (400)..(405)

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aggacaggcg tcttctactg gtgctaccag caaatgctgg aagccggaa cactgggtac 180

gttggaaacc acgtgatgtg gagtaagata aactgttagga gaaaagcatt tcgttagtgg 240

ccatgaagcc tttcaggaca tgtattgcag tatggccgg cccattacgc aattggacga 300

caacaaagac tagtattagt accacctcg ctatccacat agatcaaagc tggttaaaa 360

gagttgtgca gatgatccgt ggcacgatcg ttttttttg aattc 405

<210> 8

<211> 442

<212> DNA

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<220>

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<223> EcoRI restriction site

<220>
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<220>
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<222> (422)..(427)
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<222> (428)..(436)
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gaattattat ggagttcatg gcaactgcta tggagtttt cctactggga attggaacgg 180
tttctacgaa attaactgtc cacacgttaa aaataataat taatgcgtaa ttgttatttt 240
ttctataaca aataaaaaac tggaaatacgata cataaatttt attactttaa ttgcacttta 300
gccttagaga tattgcgttg tagtcggcgt aggtgtgtca ggggccaata tattgttccc 360
acatcggcag tgcagcacat aaactctagc gttataagaa tctatccact atcaacggtc 420
acgatcggtt tttttgaat tc 442

<210> 9
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<220>
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<220>

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<222> (101)..(195)
<223> spacer sequence

<220>
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<222> (190)..(195)
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<220>
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<222> (285)..(290)
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<220>
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gtggtgacgc atgtcgcgca agactgtaac cacgcgtctg ttgactggca ggtggtggcc      120
aatggtgatg tcagcggtga actgcgtgat gcggatcaac aggtggttgc aactggacaa      180
ggcactagcg ggatccagac gcgtggttac agtcttgcgc gacatgcgtc accacggtga      240
tatcgccac ccagggtgtc ggcgtggtgt agagcattac gctgcgatcg tcgac      295

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expression of the GUS gene (GUShp41)

<220>
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<220>
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<222> (88)..(93)
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caccacgatg ccatgttcat ctgcccagtc gac                                93

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accgccacgg ccaccgccac taccccccac tccg 94

<210> 13
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gccatatccc tacatggctg aaccacacca tcca 94

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<212> DNA
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atcatgtcac cacttccact agcagttgac aatg 94

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expression of the PDS gene (PDS42)-upper strand

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ccggtagga tcgtaatatt ccttacaagt taag 94

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